\_\_\_\_\_\_\_\_

Sequence Listing could not be accepted.

If you need help call the Patent Electronic Business Center at (866) 217-9197 (toll free).

Reviewer: Durreshwar Anjum

Timestamp: Wed Jun 06 13:00:52 EDT 2007

\*\*\*\*\*\*\*\*\*\*\*\*\*

Reviewer Comments:

Seq Id 15,16,17 has an invalid response for <213>. If <213> responses are Aritificial or Unknown please give the source of genetic material. The response mentioned is not sufficient.

\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*

## Validated By CRFValidator v 1.0.2

Application No:

10077624

Version No:

2.0

Input Set:

O

## Output Set:

**Started:** 2007-06-05 17:45:55.806

Finished: 2007-06-05 17:45:57.165

**Elapsed:** 0 hr(s) 0 min(s) 1 sec(s) 359 ms

Total Warnings: 31

Total Errors: 0

No. of SeqIDs Defined: 31

Actual SeqID Count: 31

Error code		or code	Error Description										
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	W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(6)	
	W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(7)	
	W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(8)	
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	M	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(10)	
	W.	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(11)	
	W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(12)	
	W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(13)	
	W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(14)	
	W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(15)	
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	W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(17)	
	W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(18)	
	W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(19)	
	W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(20)	

Input Set:

Output Set:

**Started:** 2007-06-05 17:45:55.806

Finished: 2007-06-05 17:45:57.165

Elapsed: 0 hr(s) 0 min(s) 1 sec(s) 359 ms

Total Warnings: 31

Total Errors: 0

No. of SeqIDs Defined: 31

Actual SeqID Count: 31

Error code Error Description

This error has occured more than 20 times, will not be displayed

## SEQUENCE LISTING

<110> THE Shi, Wenyu Morrison, Trinh, Kha Wims, Leti Chen, Li Anderson, Qi, Fengxi	Sherie m tia Maxwell
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<130> 591	57.8007.US01
<140> 100	77624
	2-02-14
<150> US	09/910,358
	1-07-19
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gccagactco	ggagaagagg ctggagtggg tcgcatccat tagtagtggt ggtacttaca 360
cctactated	agacagtgtg aagggccgat tcaccatctc cagagacaat gccaagaaca 420
ccctgtacct	gcaaatgacc agtctgaagt ctgaggacac agccatgtat tactgttcaa 480
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563

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Lys His His Ser His Arg Gly Tyr
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<211> 16
<212> PRT
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<210> 4
<211> +55
<212> 2RT
<213> Artificial sequence
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<223> Synthesized using sequential PCR techniques
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Lys His His Ser His Arg Gly Tyr Ser Gly Gly Gly Ser Gly Gly
       20
              25
Gly Gly Ser Gly Gly Gly Ser Asp Val Lys Leu Val Glu Ser Gly
      35
                       40
Gly Gly Leu Val Asn Pro Gly Gly Ser Leu Lys Leu Ser Cys Ala Ala
    50 55
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Ser Gly Phe Thr Phe Ser Ser Tyr Thr Met Ser Trp Val Arg Gln Thr 65 ' 70 Pro Glu Lys Arg Leu Glu Trp Val Ala Ser Ile Ser Ser Gly Gly Thr 90 Tyr Thr Tyr Tyr Pro Asp Ser Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ala Lys Asn Thr Leu Tyr Leu Gln Met Thr Ser Leu Lys Ser 120 115 Glu Asp Thr Ala Met Tyr Tyr Cys Ser Arg Asp Asp Gly Ser Tyr Gly . 135 130 Ser Tyr Tyr Tyr Ala Met Asp Tyr Trp Gly Gln Gly Thr Ser Val Thr 150 155 Val Ser Ser Ala Ser 165 <210> 5 <211> 533 <212> DNA <213> Artificial sequence <220> <223> Synthesized using squential PCR techniques <400> 5 ggatatccac catggacttc gggttgagct tggttttcct tgtccttact ttaaaaggtg tccagtgtaa gcggctgttt aaggagctca agttcagcct gcgcaagtac tctggtggcg 120 gtggctcggg cggaggtggg tcgggtggcg gcggatccga cgtgaagctt gtggagtctg 180 ggggaggett agtgaaccet ggagggteec tgaaactete etgtgeagee tetggattea 240 300 ctttcagtag ctataccatg tcttgggttc gccagactcc ggagaagagg ctggagtggg tcgcatccat tagtagtggt ggtacttaca cctactatcc agacagtgtg aagggccgat 360 420 tcaccatctc cagagacaat gccaagaaca ccctgtacct gcaaatgacc agtctgaagt

ctgaggacac agccatgtat tactgttcaa gagatgacgg ctcctacggc tcctattact

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480

533

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<223> Synthesized using squential PCR techniques
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<211> 155
<212> PRT
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<223> Synthesized using squential PCR techniques
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                        25
Lys Leu Val Glu Ser Gly Gly Gly Leu Val Asn Pro Gly Gly Ser Leu
      35
                     40
Lys Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Ser Tyr Thr Met
   50 55 60
Ser Trp Val Arg Gln Thr Pro Glu Lys Arg Leu Glu Trp Val Ala Ser
       70 75 80
Ile Ser Ser Gly Gly Thr Tyr Thr Tyr Tyr Pro Asp Ser Val Lys Gly
                    90
        85
Arg Phe Thr Ile Ser Arg Asp Asn Ala Lys Asn Thr Leu Tyr Leu Gln
         100 105 110
Met Thr Ser Leu Lys Ser Glu Asp Thr Ala Met Tyr Tyr Cys Ser Arg
     115 120
                               125
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<223> Primer 989

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<223> Primer 990
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cgcaagtac
<210> 13
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<223> Primer 991
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tccag
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<223> Primer 452
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<211> 18
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<223> Protegrin PG-1
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<400> 11

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5 10 15
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Gly Arg

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<211> 57

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57

<210> 17

<211> 18

<212> PRT

<213> Artificial sequence

<220>

<223> Novispirin G10

<400> 17

Lys Asn Leu Arg Arg Ile Ile Arg Lys Gly Ile His Ile Ile Lys Lys 1  $\phantom{\bigg|}$  5  $\phantom{\bigg|}$  10  $\phantom{\bigg|}$  15

Tyr Gly

<210> 18

<211> 36

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<220>

<223> Forward primer 1

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36

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<212> DNA

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<220>

<223> Reverse primer 2

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                                                                    23
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<400> 19

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<220>

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<223> Reverse primer 10
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<212> DNA
<213> Artificial sequence
<220>
<223> Forward primer 11
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gggcagcccc gagaacaac
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<211> 33
<212> DNA
<213> Artificial sequence
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<223> Reverse primer 12
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ggtggtctgc agtttacccg gggacaggga gag
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